AMENDMENTS TO THE CLAIMS:

This listing of claims will replace all prior versions, and listings, of claims in the application.

LISTING OF CLAIMS:

Claims 1 to 48 (Canceled)

49. (Currently amended) A vaccine comprising a recombinant attenuated respiratory syncytial virus, the genome of which contains the reverse complement of an mRNA coding sequence operatively linked to a polymerase binding site of a respiratory syncytial virus, and a pharmaceutically acceptable carrier, wherein the genome comprises a genetic alteration selected from (i) a deletion, (ii) an insertion, or (iii) a substitution of an entire open reading frame.

Claim 50. (Canceled)

- 51. (Previously presented) The vaccine of claim 49, wherein the genetic alteration is in a regulatory domain.
- 52. (Previously presented) The vaccine of claim 49, wherein the genetic alteration is in a functional domain.
- 53. (Previously presented) The vaccine of claim 49, wherein the virus is capable to go through only one round of replication in the host.
 - 54. (Canceled)
- 55. (Previously presented) The vaccine of claim 49, wherein the genetic alteration is an addition of one or more nucleotides.
- 56. (Previously presented) The vaccine of claim 49, wherein the genetic alteration is a deletion of one or more nucleotides.

- 57. (Currently amended) An immunogenic composition comprising a recombinant attenuated respiratory syncytial virus, the genome of which contains the reverse complement of an mRNA coding sequence operatively linked to a polymerase binding site of a respiratory syncytial virus, and a pharmaceutically acceptable carrier, wherein the genome comprises a genetic alteration selected from (i) a deletion, (ii) an insertion, or (iii) a substitution of an entire open reading frame.
- 58. (Currently amended) An immunogenic composition comprising a recombinant attenuated respiratory syncytial virus, the genome of which contains the reverse complement of an mRNA coding sequence operatively linked to a polymerase binding site of a respiratory syncytial virus, and a pharmaceutically acceptable carrier, wherein the genome comprises a modification not found in the genome of native RSV, wherein the modification is selected from (i) a deletion, (ii) an insertion, or (iii) a substitution of an entire open reading frame.

59. (Canceled)

- 60. (Previously presented) The immunogenic composition of claim 57, wherein the genetic alteration is an addition of one or more nucleotides.
- 61. (Previously presented) The immunogenic composition of claim 57, wherein the genetic alteration is a deletion of one or more nucleotides.
- 62. (Previously presented) The immunogenic composition of claim 57, wherein the genetic alteration is in a regulatory domain.
- 63. (Previously presented) The immunogenic composition of claim 57, wherein the genetic alteration is in a functional domain.

64. (Canceled)

65. (Previously presented) The immunogenic composition of claim 58, wherein the modification is an addition of one or more nucleotides.

- 66. (Previously presented) The immunogenic composition of claim 58, wherein the modification is a deletion of one or more nucleotides.
- 67. (Previously presented) The immunogenic composition of claim 58, wherein the modification is in a regulatory domain.
- 68. (Previously presented) The immunogenic composition of claim 58, wherein the modification is in a functional domain.
- 69. (Previously presented) The immunogenic composition of claim 57 or 58, wherein the virus is capable to go through only one round of replication in the host.
- 70. (Currently amended) A vaccine comprising the immunogenic composition of claim 57, 58, 59, 60, 61, 62, 63, 64, 65, 66, 67, or 68[[, or 69]].
 - 71. (New) A vaccine comprising the immunogenic composition of claim 69.
- 72. (New) The immunogenic composition of claim 57 or 58, wherein the modification affects at least one of the following the M2-2 open reading frame, the SH gene, the NS1 gene, and the NS2 gene.
- 73. (New) A pharmaceutical composition comprising (1) a recombinant attenuated respiratory syncytial virus, the genome of which contains the reverse complement of an mRNA coding sequence operatively linked to a polymerase binding site of a respiratory syncytial virus, and a pharmaceutically acceptable carrier, wherein the genome comprises a genetic alteration selected from (i) a deletion, (ii) an insertion, or (iii) a substitution of an entire open reading frame; and (2) a pharmaceutically acceptable carrier.
- 74. (New) A pharmaceutical composition comprising (1) a recombinant attenuated respiratory syncytial virus, the genome of which contains the reverse complement of an mRNA coding sequence operatively linked to a polymerase binding site of a respiratory syncytial virus, and a pharmaceutically acceptable carrier, wherein the genome comprises a modification not found in the genome of native RSV, wherein the modification is selected

- from (i) a deletion, (ii) an insertion, or (iii) a substitution of an entire open reading frame; and (2) a pharmaceutically acceptable carrier.
- 75. (New) The pharmaceutical composition of claim 73, wherein the genetic alteration is an addition of one or more nucleotides.
- 76. (New) The pharmaceutical composition of claim 732, wherein the genetic alteration is a deletion of one or more nucleotides.
- 77. (New) The pharmaceutical composition of claim 73, wherein the genetic alteration is in a regulatory domain.
- 78. (New) The pharmaceutical composition of claim 73, wherein the genetic alteration is in a functional domain.
- 79. (New) The pharmaceutical composition of claim 74, wherein the modification is an addition of one or more nucleotides.
- 80. (New) The pharmaceutical composition of claim 74, wherein the modification is a deletion of one or more nucleotides.
- 81. (New) The pharmaceutical composition of claim 74, wherein the modification is in a regulatory domain.
- 82. (New) The pharmaceutical composition of claim 74, wherein the modification is in a functional domain.
- 83. (New) The pharmaceutical composition of claim 73 or 74, wherein the virus is capable to go through only one round of replication in the host.